

Notice of References Cited	Application/Control No. 09/745,605	Applicant(s)/Patent Under Reexamination STARLING ET AL.	
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*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
	A	US-6,134,002	10-2000	Stimson, MJ. et al	
	B	US-			
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	E	US-			
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	I	US-			
	J	US-			
	K	US-			
	L	US-			
	M	US-			

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	N	WO 99/63088	12-1999		Baker k et al	
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	U	Darnel et al, Molecular Cell Biology.. New York: W H Freeman & Co; pp 248-255, 1986.
	V	Hillier et al (GenBand Accetion No. H73135 (1995))
	W	Ngo J.T, Marks J., Karplus M., Computational complexity, protein structure prediction, and the Levinthal paradox in The Protein Folding Problem, ch.14, pp. 435-508, Birkhauser, 1994.
	X	Attwood TK. Genomics. The Babel of bioinformatics. Science. 2000 Oct 20;290(5491):471-473.

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
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	V	Metzler WJ, Bajorath J, Fenderson W, Shaw SY, Constantine KL, Naemura J, Leytze G, Peach RJ, Lavoie TB, Mueller L, Linsley PS. Solution structure of human CTLA-4 and delineation of a CD80/CD86 binding site conserved in CD28. Nat Struct Biol. 1997 Jul;4(7):
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